\_\_\_\_\_\_

Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Fri Oct 05 11:45:28 EDT 2007

\_\_\_\_\_\_

## Validated By CRFValidator v 1.0.3

Application No: 10578840 Version No: 1.0

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Input Set:

Output Set:

**Started:** 2007-09-21 18:26:57.594 **Finished:** 2007-09-21 18:27:00.392

**Elapsed:** 0 hr(s) 0 min(s) 2 sec(s) 798 ms

Total Warnings: 121

No. of SeqIDs Defined: 121

Actual SeqID Count: 121

Total Errors:

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Input Set:

Output Set:

**Started:** 2007-09-21 18:26:57.594 **Finished:** 2007-09-21 18:27:00.392

**Elapsed:** 0 hr(s) 0 min(s) 2 sec(s) 798 ms

Total Warnings: 121
Total Errors: 0
No. of SeqIDs Defined: 121

Actual SeqID Count: 121

Error code Error Description

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	TSUCHIYA, MASAYUKI	
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gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag
                                                                   96
Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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                                                                   144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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Ala Asn His Val Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu
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gtc cac tcc cag gtg cag ctg gtg cag tct ggg gct gag gtg aag aag
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Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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Glu Lys Phe Lys Asp Lys Val Thr Met Thr Ser Asp Thr Ser Thr Ser
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aca gtc tac atg gag ttg agc agt ctc aga tct gag gac acg gcc gtc
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Thr Val Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val
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tat tat tgt gct aga ggg ggt tac tat act tac gac gac tgg ggc caa
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Tyr Tyr Cys Ala Arg Gly Gly Tyr Tyr Thr Tyr Asp Asp Trp Gly Gln
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Val His Ser Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys
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cct ggg gcc tca gtg aag gtt tcc tgt aag gca tct gga tac acc ttc
                                                                    144
Pro Gly Ala Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe
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<213> A